

10536705\_70439-00003\_ST25.txt  
SEQUENCE LISTING

<110> Montelione, Gaetano  
Das, Kalyan  
Arnold, Eddy

<120> Ribosomal RNA Methyltransferases Rima: Target Validation and  
Processes for Developing an Inhibitor Assay and Identification of  
Candidate Inhibitors

<130> 70439.00003

<140> 10/536,705  
<141> 2006-03-08

<150> PCT/US04/20244  
<151> 2004-06-26

<150> US 60/482,722  
<151> 2003-06-27

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<170> PatentIn version 3.4

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<212> PRT  
<213> Escherichia coli

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Asn Ser Tyr Ile Cys Pro Gln Arg His Gln Phe Asp Met Ala Lys Glu  
20 25 30

Gly Tyr Val Asn Leu Leu Pro Val Gln His Lys Arg Ser Arg Asp Pro  
35 40 45

Gly Asp Ser Ala Glu Met Met Gln Ala Arg Arg Ala Phe Leu Asp Ala  
50 55 60

Gly His Tyr Gln Pro Leu Arg Asp Ala Ile Val Ala Gln Leu Arg Glu  
65 70 75 80

Arg Leu Asp Asp Lys Ala Thr Ala Val Leu Asp Ile Gly Cys Gly Glu  
85 90 95

Gly Tyr Tyr Thr His Ala Phe Ala Asp Ala Leu Pro Glu Ile Thr Thr  
100 105 110

Phe Gly Leu Asp Val Ser Lys Val Ala Ile Lys Ala Ala Lys Arg  
115 120 125

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Tyr Pro Gln Val Thr Phe Cys Val Ala Ser Ser His Arg Leu Pro Phe  
130 135 140

Ser Asp Thr Ser Met Asp Ala Ile Ile Arg Ile Tyr Ala Pro Cys Lys  
145 150 155 160

Ala Glu Glu Leu Ala Arg Val Val Lys Pro Gly Gly Trp Val Ile Thr  
165 170 175

Ala Thr Pro Gly Pro Arg His Leu Met Glu Leu Lys Gly Leu Ile Tyr  
180 185 190

Asn Glu Val His Leu His Ala Pro His Ala Glu Gln Leu Glu Gly Phe  
195 200 205

Thr Leu Gln Gln Ser Ala Glu Leu Cys Tyr Pro Met Arg Leu Arg Gly  
210 215 220

Asp Glu Ala Val Ala Leu Leu Gln Met Thr Pro Phe Ala Trp Arg Ala  
225 230 235 240

Lys Pro Glu Val Trp Gln Thr Leu Ala Ala Lys Glu Val Phe Asp Cys  
245 250 255

Gln Thr Asp Phe Asn Ile His Leu Trp Gln Arg Ser Tyr  
260 265

<210> 2  
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<212> PRT  
<213> Salmonella typhimurium

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Asn Ser Val Ile Cys Pro Gln Arg His Gln Phe Asp Val Ala Lys Glu  
20 25 30

Gly Tyr Ile Asn Leu Leu Pro Val Gln His Lys Arg Ser Arg Asp Pro  
35 40 45

Gly Asp Ser Ala Glu Met Met Gln Ala Arg Arg Ala Phe Leu Asp Ala  
50 55 60

Gly His Tyr Gln Pro Leu Arg Asp Ala Val Ile Asn Leu Leu Arg Glu  
65 70 75 80

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Arg Leu Asp Gln Ser Ala Thr Ala Ile Leu Asp Ile Gly Cys Gly Glu  
85 90 95

Gly Tyr Tyr Thr His Ala Phe Ala Glu Ala Leu Pro Gly Val Thr Thr  
100 105 110

Phe Gly Leu Asp Val Ala Lys Thr Ala Ile Lys Ala Ala Lys Arg  
115 120 125

Tyr Ser Gln Val Lys Phe Cys Val Ala Ser Ser His Arg Leu Pro Phe  
130 135 140

Ala Asp Ala Ser Met Asp Ala Val Ile Arg Ile Tyr Ala Pro Cys Lys  
145 150 155 160

Ala Gln Glu Leu Ala Arg Val Val Lys Pro Gly Gly Trp Val Val Thr  
165 170 175

Ala Thr Pro Gly Pro His His Leu Met Glu Leu Lys Gly Leu Ile Tyr  
180 185 190

Asp Glu Val Arg Leu His Ala Pro Tyr Thr Glu Gln Leu Asp Gly Phe  
195 200 205

Thr Leu Gln Gln Ser Thr Arg Leu Ala Tyr His Met Gln Leu Thr Ala  
210 215 220

Glu Ala Ala Val Ala Leu Leu Gln Met Thr Pro Phe Ala Trp Arg Ala  
225 230 235 240

Arg Pro Asp Val Trp Glu Gln Leu Ala Ala Ser Ala Gly Leu Ser Cys  
245 250 255

Gln Thr Asp Phe Asn Leu His Leu Trp Gln Arg Asn Arg  
260 265

<210> 3  
<211> 279  
<212> PRT  
<213> Yersinia pestis

<400> 3

Met Ser Tyr Gln Cys Pro Leu Cys His Gln Ala Leu Gln Leu Ser Gln  
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Gln Gln Trp Cys Cys Ser Asn Asn His Gln Phe Asp Cys Ala Lys Glu  
20 25 30

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Gly Tyr Val Asn Leu Met Pro Val Gln His Lys Gly Ser Lys Gln Pro  
35 40 45

Gly Asp Ser Pro Glu Met Met Gln Ala Arg Arg Ala Phe Leu Asp Ala  
50 55 60

Gly Tyr Tyr Gln Pro Leu Gln Gln Arg Val Ser Glu Ile Leu Asp Glu  
65 70 75 80

Ala Leu Pro Leu Asp Ala Thr Arg Leu Leu Asp Ile Gly Cys Gly Glu  
85 90 95

Gly Tyr Tyr Thr Ala Ala Val Ala Asp Arg Leu Asn Lys Leu Arg Gln  
100 105 110

Met Ala Ile Phe Gly Leu Asp Val Ala Lys Val Ala Val Arg Tyr Gly  
115 120 125

Ala Lys Arg Tyr His Gln Val Asn Phe Cys Val Ala Ser Ser His Arg  
130 135 140

Leu Pro Phe Ala Asn Gly Ala Leu Asp Ala Val Leu Arg Ile Tyr Ala  
145 150 155 160

Pro Cys Lys Ala Val Glu Leu Ala Arg Thr Val Lys Pro Gly Gly Ile  
165 170 175

Val Val Thr Val Ala Pro Gly Pro Arg His Leu Tyr Gln Leu Lys Ala  
180 185 190

Leu Ile Tyr Ala Gln Val Gln Leu His Asp Asp Thr Glu Glu His Leu  
195 200 205

Asp Gly Phe Glu Leu Ile Arg Arg Glu Thr Leu Ala Tyr Asp Met Lys  
210 215 220

Leu Thr Gly Glu Gln Gly Phe Asn Leu Leu Gln Met Thr Pro Phe Ala  
225 230 235 240

Trp Arg Ala Ser Val Asp Thr Gly Gln Lys Leu Ala Ala Gln Pro Ser  
245 250 255

Phe Ser Cys Glu Thr Asp Phe Val Ile Ser Leu His Arg Arg Lys Thr  
260 265 270

Asp Asn Pro Gln Asn Asp Ile  
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<213> Vibrio cholerae

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Phe Asp Val Ala Lys Glu Gly Tyr Val Asn Leu Met Pro Val Gln His
 35                      40                      45

Lys Arg Ser Lys Asp Pro Gly Asp Asn Lys Glu Met Thr Gln Ala Arg
 50                      55                      60

Arg Arg Phe Leu His Thr Gly His Tyr Ala Pro Met Arg Glu Lys Val
 65                      70                      75                      80

Ala Thr Leu Cys Gln Thr Tyr Leu Thr Gly Arg Gln Gln Thr Leu Leu
          85                      90                      95

Asp Ile Gly Cys Gly Glu Gly Tyr Tyr Thr Asp Phe Phe Ala Lys Ala
      100                      105                      110

Leu Ser Gln Gln Asp Ser Glu Ala Gln Ile Leu Gly Leu Asp Ile Ser
      115                      120                      125

Lys Ile Ala Ile Arg Tyr Ala Ala Lys Arg Tyr Pro Glu Cys Gln Phe
      130                      135                      140

Ala Val Ala Ser Ser His Arg Leu Pro Phe Ala Asn Gln Ser Leu Asp
      145                      150                      155                      160

Gly Val Ile Arg Ile Tyr Ala Pro Cys Lys Asp Thr Glu Leu Glu Arg
      165                      170                      175

Cys Ile Lys Ile Gly Gly Ile Val Ile Thr Val Thr Pro Ala Ala Arg
      180                      185                      190

His Leu Tyr Gln Phe Lys Gln Gly Ile Tyr Asp Gln Val Arg Leu His
      195                      200                      205

Glu Glu Gln Pro Glu Thr Leu Ser Gly Phe Glu Leu Val Glu Glu Cys
      210                      215                      220

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Lys Leu His Tyr Pro Met Ala Leu Asn Gly Ser Glu Ala Ala Asp Leu  
225 230 235 240

Leu Gln Met Thr Pro Phe Ala Trp Arg Ala Ser Glu Asp Phe Lys His  
245 250 255

Arg Val Ser Gln Ser Asp Thr Phe Glu Cys Glu Ala Asp Phe Met Leu  
260 265 270

Arg Val Tyr Arg Arg Lys  
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<212> PRT  
<213> Pseudomonas putida

<400> 5

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Gly Val Val Cys Pro Ala Gly His Arg Phe Asp Arg Ala Arg Gln Gly  
20 25 30

Tyr Leu Asn Leu Leu Pro Val Gln His Lys Asn Ser Arg Asp Pro Gly  
35 40 45

Asp Asn Gln Ala Met Val Glu Ala Arg Arg Asp Phe Leu Asp Ala Gly  
50 55 60

His Tyr Ala Pro Val Ala Arg Arg Leu Ala Glu Leu Ala Ala Glu Arg  
65 70 75 80

Gln Pro Gly Ala Trp Leu Asp Ile Gly Cys Gly Glu Gly Tyr Tyr Thr  
85 90 95

Ala Gln Ile Ala Gln Ala Leu Pro Ala Ala Asp Gly Tyr Ala Leu Asp  
100 105 110

Ile Ser Arg Glu Ala Val Lys Arg Ala Cys Arg Arg Ala Ser Ala Val  
115 120 125

Thr Trp Met Val Ala Ser Met Ala Arg Val Pro Leu Thr Asp Ala Ser  
130 135 140

Cys Gln Phe Ile Ala Ser Val Phe Ser Pro Leu Asp Trp Ala Glu Ala  
145 150 155 160

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Lys Arg Leu Leu Ser Pro Gly Gly Gly Leu Met Arg Val Gly Pro Thr  
165 170 175

Ser Gly His Leu Met Glu Leu Arg Glu Val Leu Tyr Asp Glu Val Arg  
180 185 190

Pro Tyr Ala Asp Asp Lys His Leu Ala Leu Val Pro Glu Gly Met Ala  
195 200 205

His Ala His Ser Glu Thr Leu Glu Phe Arg Leu Ser Leu Ala Ala Pro  
210 215 220

Lys Ala Arg Ala Asp Leu Leu Ala Met Thr Pro His Gly Trp Arg Ala  
225 230 235 240

Ser Ala Glu Lys Arg Ala Arg Val Ile Asp Gln Pro Glu Pro Phe Glu  
245 250 255

Val Thr Val Ser Met Arg Tyr Asp Tyr Phe Val Arg Gln Asp  
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<210> 6  
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<213> Pseudomonas aeruginosa  
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Gly Val Ala Cys Pro Ala Gly His Arg Phe Asp Arg Ala Arg Gln Gly  
20 25 30

Tyr Leu Asn Leu Leu Pro Val Gln His Lys Lys Ser Leu Asp Pro Gly  
35 40 45

Asp Asn Ala Ala Met Val Glu Ala Arg Arg Gln Phe Leu Gly Ala Gly  
50 55 60

His Tyr Ala Pro Leu Ala Arg Arg Leu Ala Glu Leu Ala Ala Glu Arg  
65 70 75 80

Ala Pro Arg Arg Trp Leu Asp Ile Gly Cys Gly Glu Gly Tyr Tyr Ser  
85 90 95

Ala Gln Leu Gly Glu Ala Leu Gly Asp Ala Glu Gly Tyr Ala Leu Asp  
100 105 110

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Ile Ser Arg Glu Ala Val Lys Arg Ala Cys Arg Arg Ala Pro Gln Leu  
115 120 125

Thr Trp Leu Val Ala Ser Met Ala Arg Leu Pro Leu Ala Glu Ala Ser  
130 135 140

Cys Glu Leu Ile Ala Ser Val Phe Ser Pro Ile Asp Trp Asn Glu Ala  
145 150 155 160

Val Arg Val Leu Ala Pro Gly Gly Gly Val Leu Arg Leu Gly Pro Ala  
165 170 175

Ser Ala His Leu Leu Glu Leu Arg Gln Arg Leu Tyr Asp Asp Val Arg  
180 185 190

Asp Tyr Ala Asp Asp Lys His Leu Ala Gly Leu Pro Ala Pro Leu Ser  
195 200 205

Leu Arg His Thr Glu Thr Leu Ser Phe Arg Leu Ala Leu Asp Ser Tyr  
210 215 220

Glu Ala Arg Glu Asn Leu Leu Ala Met Thr Pro His Gly Trp Arg Val  
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Asn Ala Glu Arg Arg Ala Arg Ile Leu Ala Glu Pro Phe Glu Val Ser  
245 250 255

Val Ala Val Arg Tyr Asp Trp Leu Gln Arg Asp  
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<210> 7  
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<213> Streptomyces fradiae

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Met Arg Lys Asn Val Val Arg Tyr Leu Arg Cys Pro His Cys Ala Ala  
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Pro Leu Arg Ser Ser Asp Arg Thr Leu Arg Cys Glu Asn Gly His Thr  
20 25 30

Phe Asp Val Ala Arg Gln Gly Tyr Val Asn Leu Leu Arg Arg Pro Thr  
35 40 45

Lys Leu Ala Ala Asp Thr Thr Asp Met Val Ala Ala Arg Ala Ala Leu  
50 55 60



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Leu Asp Ser Gly His Tyr Ala Pro Leu Thr Glu Arg Leu Ala Gly Thr  
 65 70 75 80  
 Ala Arg Arg Ala Ala Gly Ala Gly Ala Pro Asp Cys Val Val Asp Ile  
 85 90 95  
 Gly Gly Gly Thr Gly His His Leu Ala Arg Val Leu Glu Glu Phe Glu  
 100 105 110  
 Asp Ala Glu Gly Leu Leu Leu Asp Met Ser Lys Pro Ala Val Arg Arg  
 115 120 125  
 Ala Ala Arg Ala His Pro Arg Ala Ser Ser Ala Val Ala Asp Val Trp  
 130 135 140  
 Asp Thr Leu Pro Leu Arg Asp Gly Ala Ala Ala Met Ala Leu Asn Val  
 145 150 155 160  
 Phe Ala Pro Arg Asn Pro Pro Glu Ile Arg Arg Ile Leu Arg Pro Gly  
 165 170 175  
 Gly Thr Leu Leu Val Val Thr Pro Gln Gln Asp His Leu Ala Glu Leu  
 180 185 190  
 Val Asp Ala Leu Gly Leu Leu Arg Val Arg Asp His Lys Glu Gly Arg  
 195 200 205  
 Leu Ala Glu Gln Leu Ala Pro His Phe Glu Ala Val Gly Gln Glu Arg  
 210 215 220  
 Leu Arg Thr Thr Leu Arg Leu Asp His Asp Ala Leu Gly Arg Val Val  
 225 230 235 240  
 Ala Met Gly Pro Ser Ser Trp His Gln Asp Pro Asp Glu Leu Ala Arg  
 245 250 255  
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 Phe Thr Val Cys Arg Pro Leu Pro  
 275 280

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 35 40 45  
 Lys Pro Val Lys Thr Ser Tyr Gly Ala Glu Leu Phe Glu Ala Arg Ser  
 50 55 60  
 Arg Leu Ile Gly Glu Cys Gly Phe Phe Asp Pro Leu His Asp Ala Ile  
 65 70 75 80  
 Ala Glu Leu Ile Ser His Pro Lys Ser Gly His Glu Ala Phe Thr Ile  
 85 90 95  
 Leu Asp Ser Gly Cys Gly Glu Gly Ser His Leu Asn Ala Leu Cys Gly  
 100 105 110  
 Phe Asp Tyr Ala Gly Lys Ala Ala Ile Gly Thr Gly Ile Asp Leu Ser  
 115 120 125  
 Lys Asp Gly Ile Leu Lys Ala Ser Lys Ala Phe Lys Asp Leu Met Trp  
 130 135 140  
 Ala Val Ala Asp Val Ala Arg Ala Pro Phe His Asp Arg Gln Phe Asp  
 145 150 155 160  
 Val Val Leu Ser Ile Phe Ser Pro Ser Asn Tyr Ala Glu Phe His Arg  
 165 170 175  
 Leu Leu Lys Asn Asp Gly Met Leu Ile Lys Val Val Pro Arg Ser Asp  
 180 185 190  
 Tyr Leu Ile Glu Leu Arg Gln Phe Leu Tyr Thr Asp Ser Pro Arg Arg  
 195 200 205  
 Thr Tyr Ser Asn Thr Ala Ala Val Glu Arg Phe Thr Ala Asn Ala Ala  
 210 215 220  
 His Ser Arg Pro Val Arg Leu Arg Tyr Val Lys Thr Leu Asp Gln Gln  
 225 230 235 240  
 Ala Ile His Trp Leu Leu Lys Met Thr Pro Leu Ala Trp Ser Ala Pro

Lys Asp Arg Val Ser Leu Leu Lys Glu Met Lys Ser Ala Asp Ile Thr  
 260 265 270

Val Asp Val Asp Ile Leu Ile Gly Met Lys  
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<210> 9  
 <211> 273  
 <212> PRT  
 <213> Lactococcus lactis

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Glu Met Leu Arg Cys Pro Ile Cys His Gly Lys Phe Gln Leu Ala Ala  
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Tyr Ala Leu Lys Cys Glu Asn Asn His Thr Tyr Asn Leu Asn Lys Lys  
 35 40 45

Gly Tyr Val Asn Phe Leu Gln Thr Lys Ala Asp Thr Glu His Tyr Thr  
 50 55 60

Arg Lys Met Phe Glu Pro Arg Arg Arg Leu Ile Gln Ala Gly Met Tyr  
 65 70 75 80

Gln Asn Leu Leu Thr Glu Ile Gln Lys Ser Phe Val Ser Gly Asn Leu  
 85 90 95

Leu Asp Val Gly Thr Gly Glu Gly Ser Phe Leu Glu Leu Leu Glu Gly  
 100 105 110

Ala Gly Ala Lys Phe Ala Phe Asp Ile Ala Lys Asp Gly Ile Glu Met  
 115 120 125

Ala Thr Glu Leu Asp Thr Glu Ser Phe Leu Ser Leu Ala Asp Leu Thr  
 130 135 140

Asn Leu Pro Phe Ala Asp Glu Ser Leu Ser Val Ile Leu Asn Ile Phe  
 145 150 155 160

Thr Pro Ser Asn Tyr Ala Glu Phe His Arg Val Leu Thr Glu Asn Gly  
 165 170 175

Arg Val Ile Lys Ile Ile Pro Asp Arg Asn Tyr Leu His Glu Leu Arg  
 Page 11

Glu Val Tyr Gln Leu Pro Val Asp Tyr Asp Asn Gln Ala Val Ile Glu  
195 200 205

Arg Phe Lys Glu Glu Phe Pro Lys Asn Thr Gln Gln Thr Ile Asp Tyr  
210 215 220

Thr Phe Glu Ile Pro Glu Asn Leu Arg Gln Asp Phe Leu Leu Met Ser  
225 230 235 240

Pro Leu Glu Trp Ser Val Ser Glu Glu Arg Lys Lys Phe Ala Lys Glu  
245 250 255

Asn Pro Pro Lys Thr Ala Arg Ile His Val Gln Ile Leu Ile Gly Ile  
260 265 270

Lys

<210> 10

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<212> PRT

<213> Streptococcus pneumoniae

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Asn Phe Lys Cys Cys Asn Arg His Ser Phe Asp Leu Ala Lys Phe Gly  
35 40 45

Tyr Val Asn Leu Val Pro Gln Ile Lys Gln Ser Ala Asn Tyr Asp Lys  
50 55 60

Glu Asn Phe Gln Asn Arg Gln Gln Ile Leu Glu Ala Gly Phe Tyr Gln  
65 70 75 80

Ala Ile Leu Asp Ala Val Ser Asp Leu Leu Ala Ser Ser Lys Thr Thr  
85 90 95

Thr Thr Ile Leu Asp Ile Gly Cys Gly Glu Gly Phe Tyr Ser Arg Lys  
100 105 110

Leu Gln Glu Ser His Ser Glu Lys Thr Phe Tyr Ala Phe Asp Ile Ser

115

Lys Asp Ser Val Gln Ile Ala Ala Lys Ser Glu Pro Asn Trp Ala Val  
130 135

Asn Trp Phe Val Gly Asp Leu Ala Arg Leu Pro Ile Lys Asp Ala Asn  
145 150 155 160

Met Asp Ile Leu Leu Asp Ile Phe Ser Pro Ala Asn Tyr Gly Glu Phe  
165 170 175

Arg Arg Val Leu Ser Lys Asp Gly Ile Leu Ile Lys Val Ile Pro Thr  
180 185 190

Glu Asn His Leu Lys Glu Ile Arg Gln Arg Val Gln Asp Gln Leu Thr  
195 200 205

Asn Lys Glu Tyr Ser Asn Gln Asp Ile Lys Glu His Phe Gln Glu His  
210 215 220

Phe Thr Ile Leu Ser Ser Gln Thr Ala Ser Leu Thr Lys Thr Ile Thr  
225 230 235 240

Ala Glu Gln Leu Gln Ala Leu Leu Ser Met Thr Pro Leu Leu Phe His  
245 250 255

Val Asp Gln Ser Lys Ile Asp Trp Ser Gln Leu Thr Glu Ile Thr Ile  
260 265 270

Glu Ala Glu Ile Leu Val Gly Lys Ala Phe  
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<223> E. coli R1mA fragment

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<210> 13

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<213> Artificial

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<223> E. coli R1mA fragment

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<210> 14

<211> 6

<212> PRT

<213> Artificial

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<223> E. coli R1mA fragment

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Leu Leu Gln Met Thr Pro  
1 5

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<212> PRT

<213> Artificial

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<223> E. coli R1mA fragment

<400> 15

Leu Asp Val Ser Lys  
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<210> 16

<211> 3

<212> PRT

<213> Artificial

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<223> E. coli R1mA fragment

<400> 16

Met Thr Pro  
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 <211> 4  
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<210> 18  
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<220>  
 <223> E. coli R1mA fragment

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&lt;223&gt; E. coli R1mA methyltransferase domain

&lt;400&gt; 21

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Leu Arg Asp Ala Ile Val Ala Gln Leu Arg Glu Arg Leu Asp Asp Lys  
 20 25 30

Ala Thr Ala Val Leu Asp Ile Gly Cys Gly Glu Gly Tyr Tyr Thr His  
 35 40 45

Ala Phe Ala Asp Ala Leu Pro Glu Ile Thr Thr Phe Gly Leu Asp Val  
 50 55 60

Ser Lys Val Ala Ile Lys Ala Ala Ala Lys Arg Tyr Pro Gln Val Thr  
 65 70 75 80

Phe Cys Val Ala Ser Ser His Arg Leu Pro Phe Ser Asp Thr Ser Met  
 85 90 95

Asp Ala Ile Ile Arg Ile Tyr Ala Pro Cys Lys Ala Glu Glu Leu Ala  
 100 105 110

Arg Val Val Lys Pro Gly Gly Trp Val Ile Thr Ala Thr Pro Gly Pro  
 115 120 125

Arg His Leu Met Glu Leu Lys Gly Leu Ile Tyr Asn Glu Val His Leu  
 130 135 140

His Ala Pro His Ala Glu Gln Leu Glu Gly Phe Thr Leu Gln Gln Ser  
 145 150 155 160

Ala Glu Leu Cys Tyr Pro Met Arg Leu Arg Gly Asp Glu Ala Val Ala  
 165 170 175

Leu Leu Gln Met Thr Pro Phe Ala Trp Arg Ala Lys Pro Glu Val Trp  
 180 185 190

Gln Thr Leu Ala Ala Lys Glu Val Phe Asp Cys Gln Thr Asp Phe Asn  
 195 200 205

Ile His Leu Trp Gln Arg Ser Tyr  
 210 215

&lt;210&gt; 22

&lt;211&gt; 12

&lt;212&gt; PRT



<213> Artificial

<220>

<223> E. coli R1mA linker

<400> 22

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<210> 23

<211> 37

<212> PRT

<213> Artificial

<220>

<223> E. coli R1mA Zn-binding domain

<400> 23

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20 25 30

Gly Tyr Val Asn Leu  
35